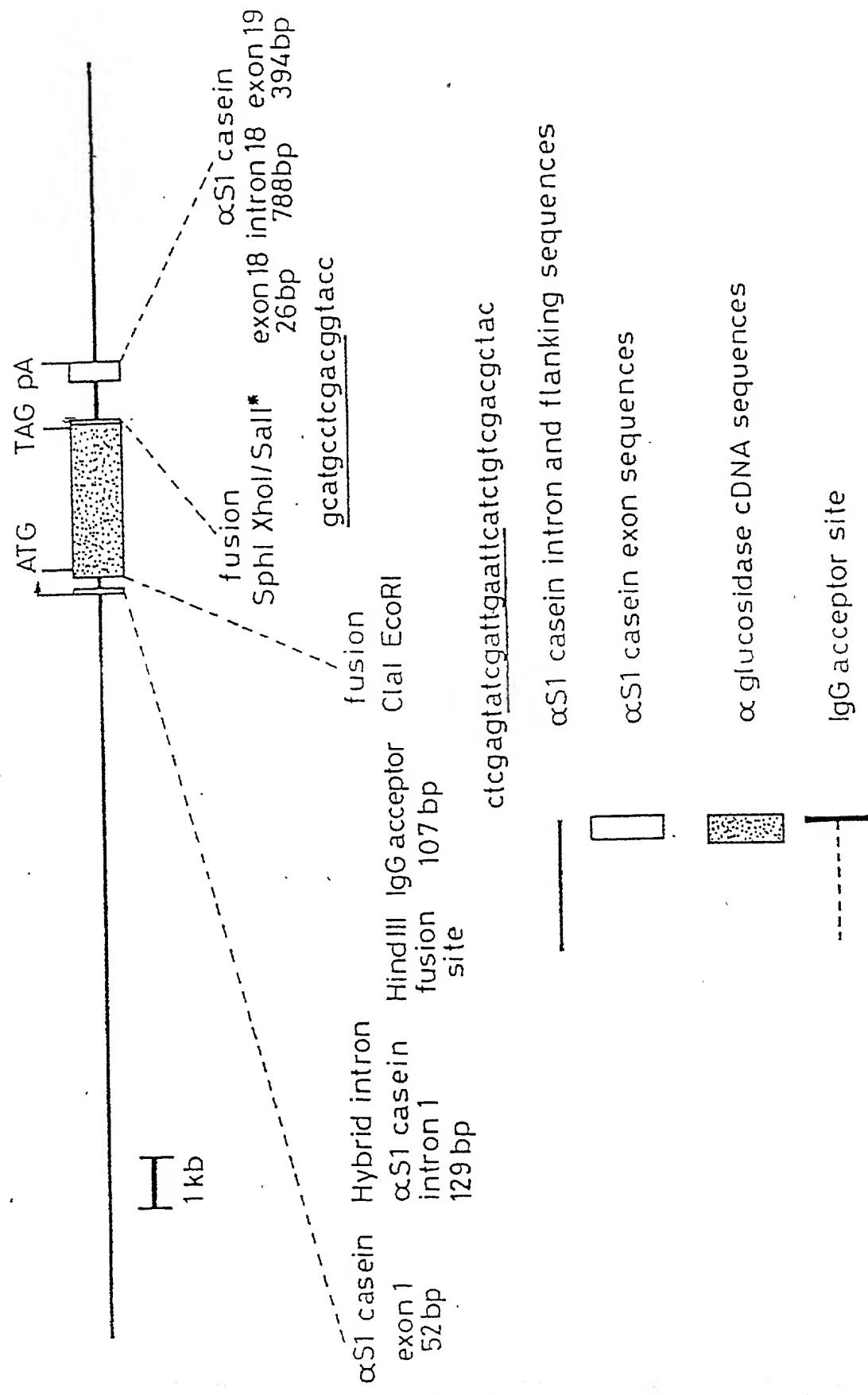


Fig. 1.



α -glucosidase constructs

Fig. 2.A

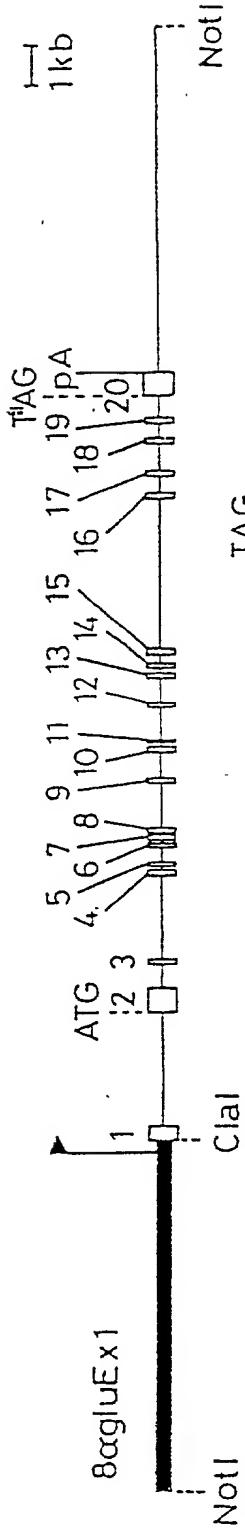


Fig. 2.B.

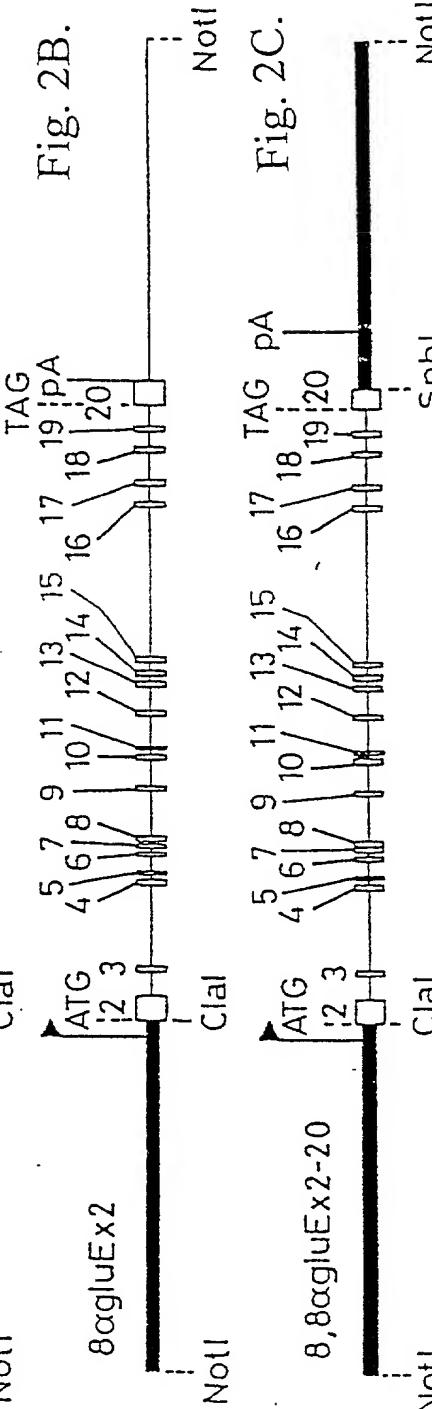
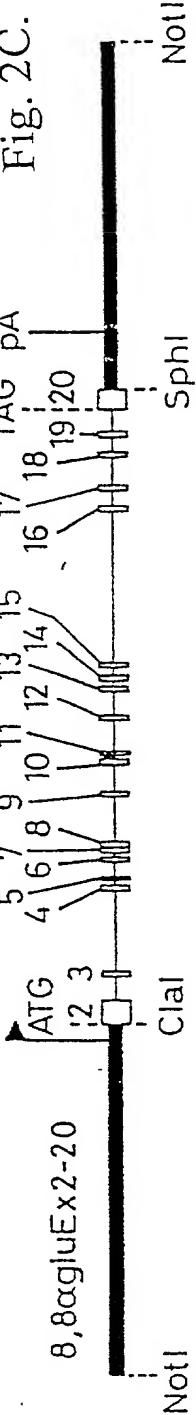


Fig. 2.C.



Transcription Initiation site.

■ α s₁ casein sequence, promoter or 3' untranslated region.

2 3 The boxes represent the exons in the α -glucosidase sequence, the thin line represents the intron sequences.

The numbers above the boxes are the exon numbers

PA = polyadenylation signal.

ATG = translation initiation site.

TAG = translation stop codon

Fig. 3A.

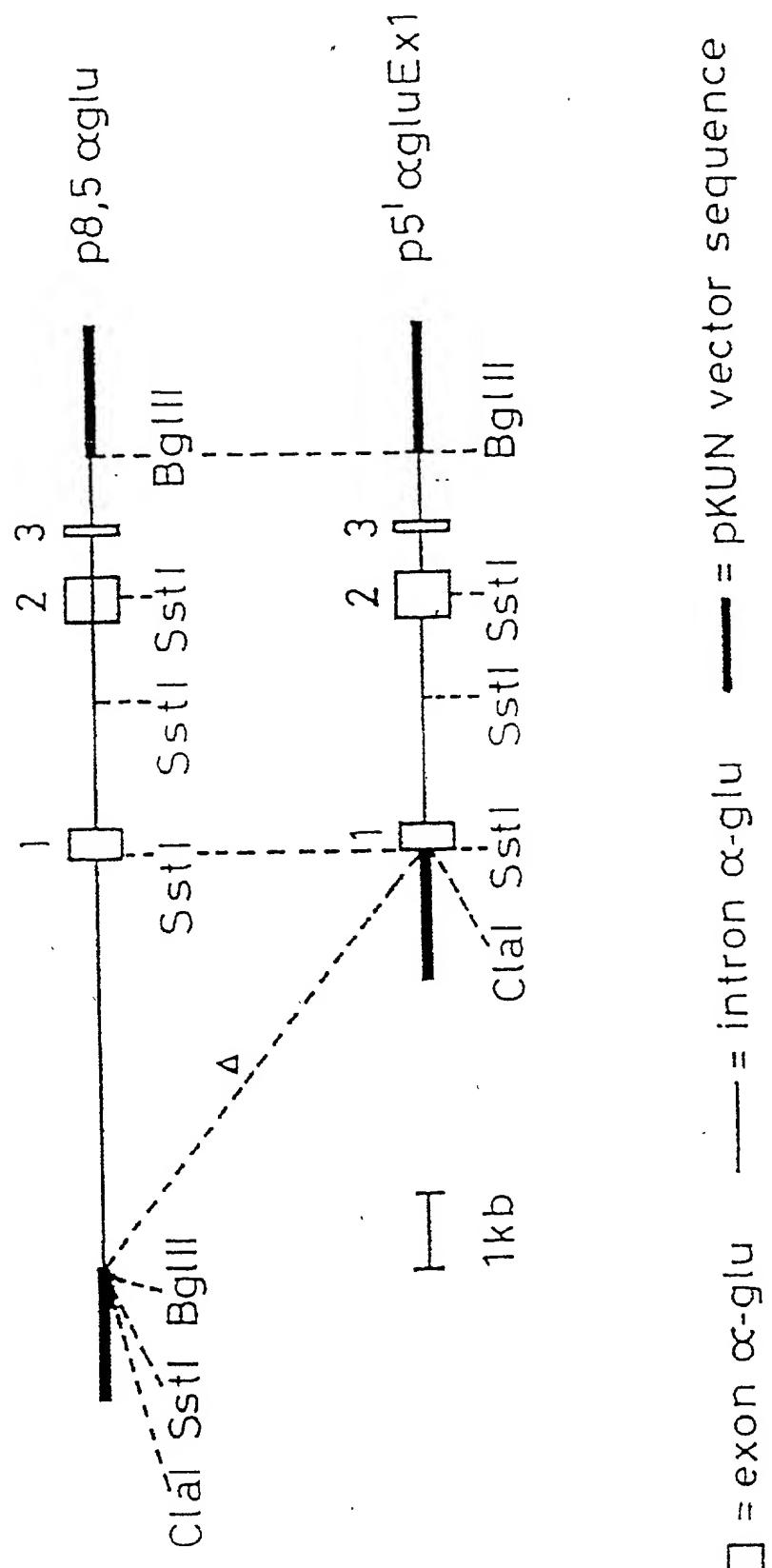


Fig. 3B.

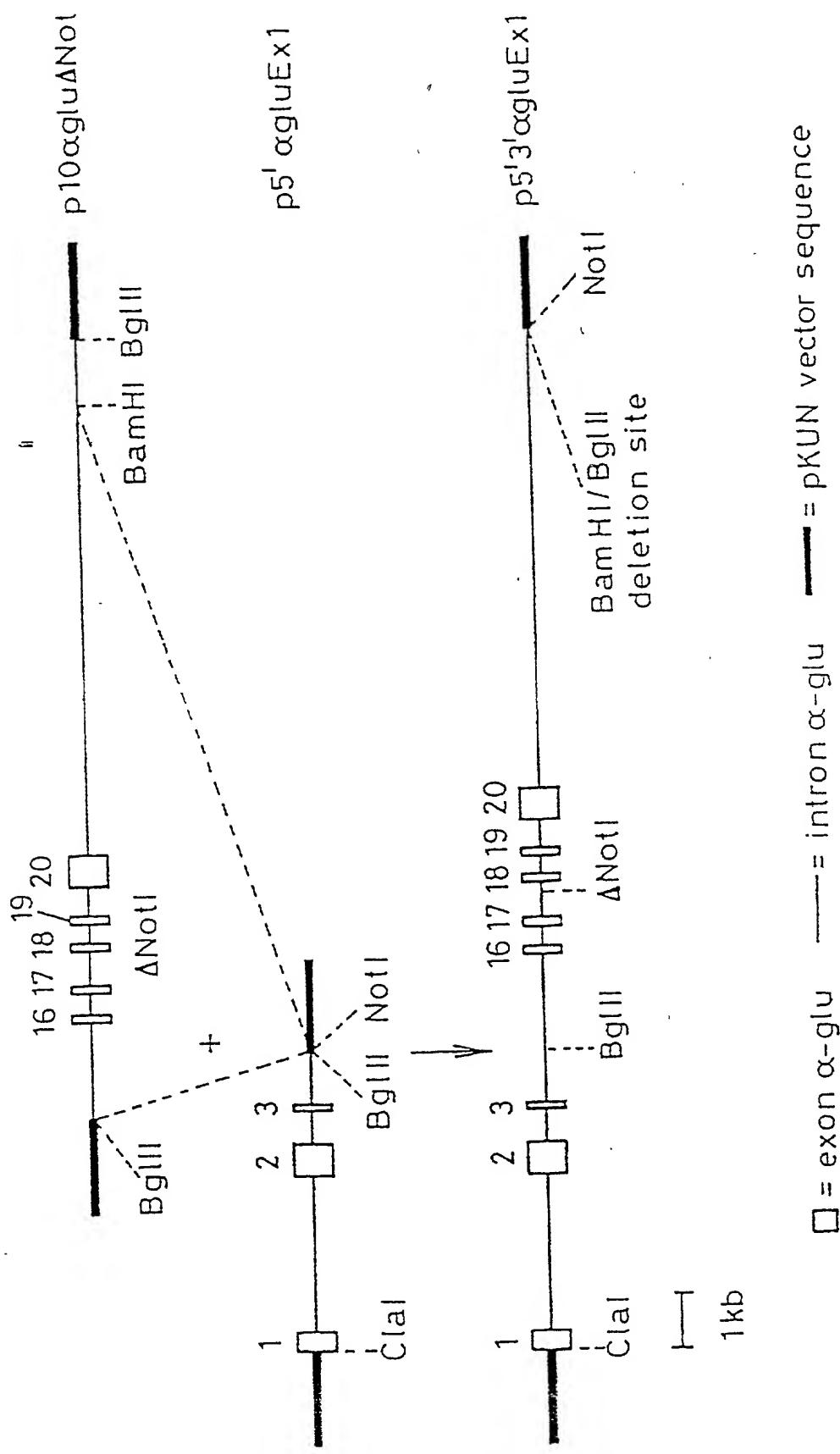


Fig. 3.C.

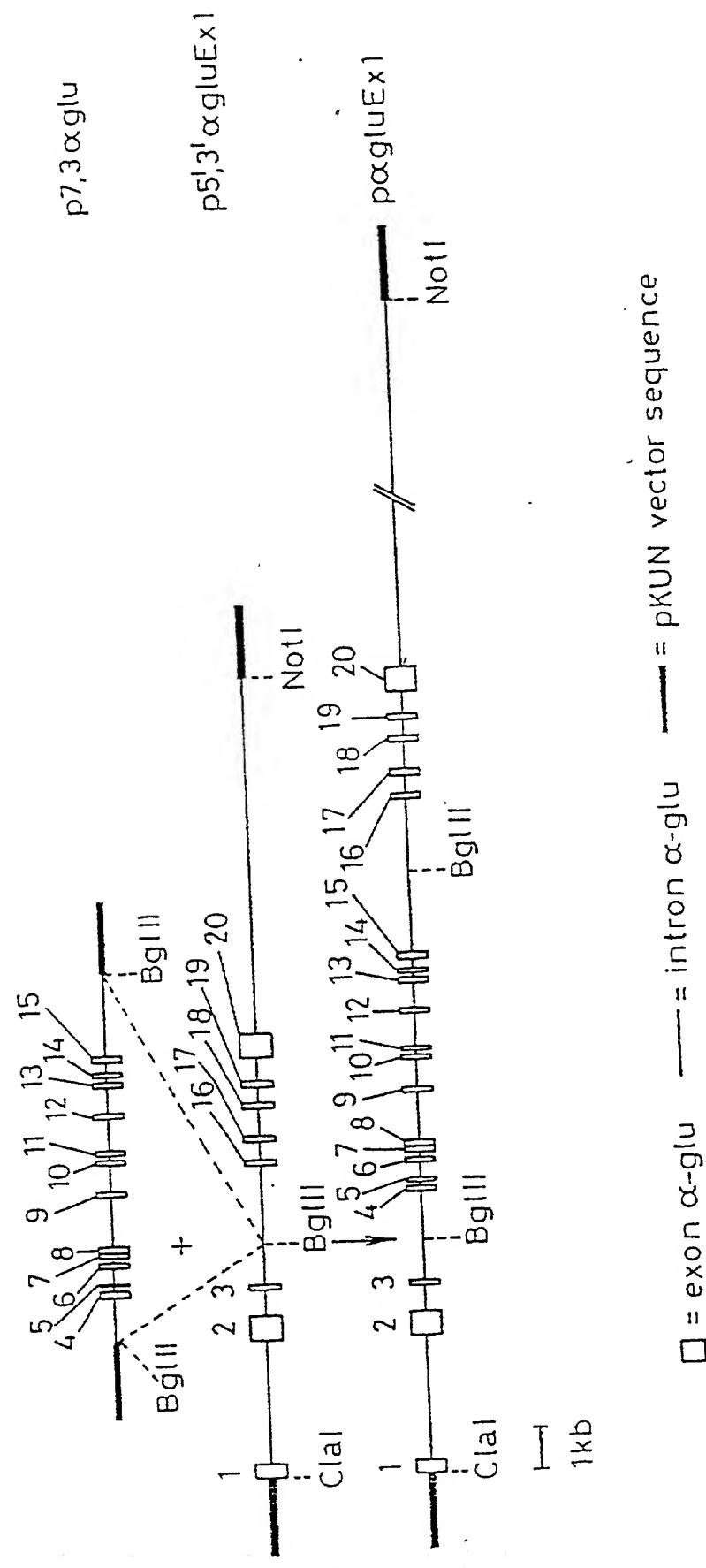


Fig. 4. A.

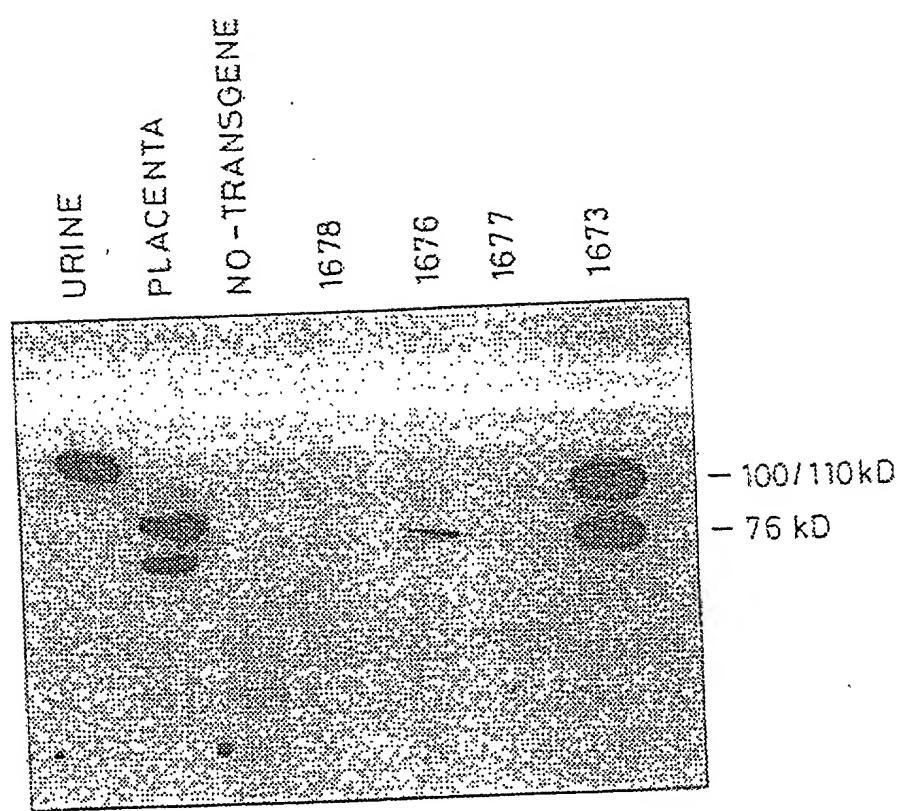


Fig. 4. B.

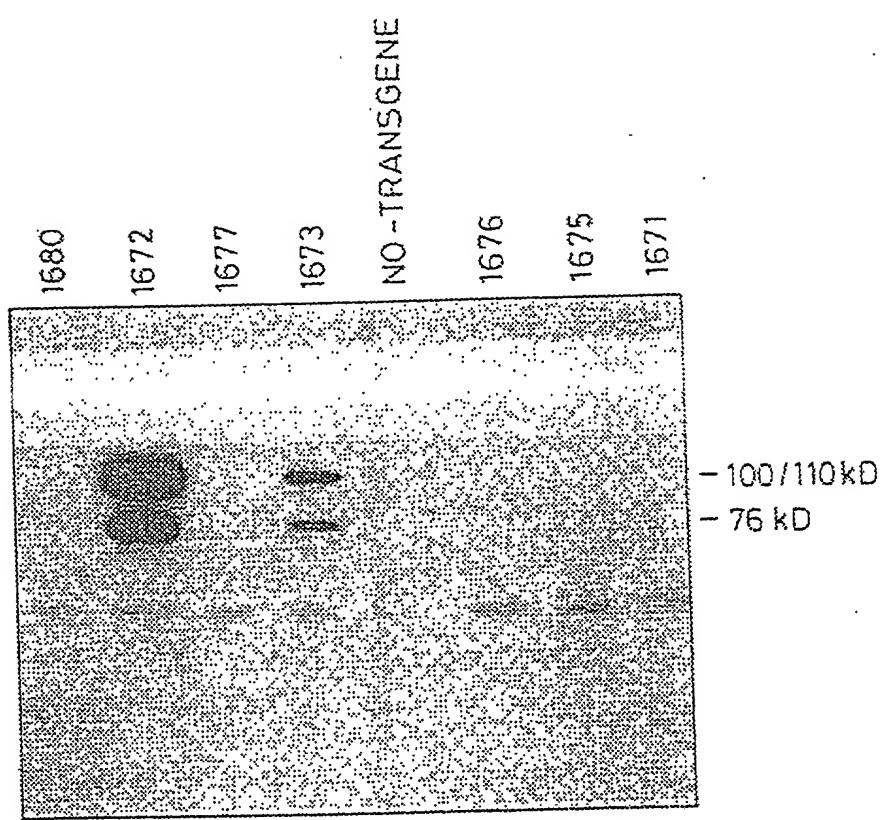


Fig. 5.

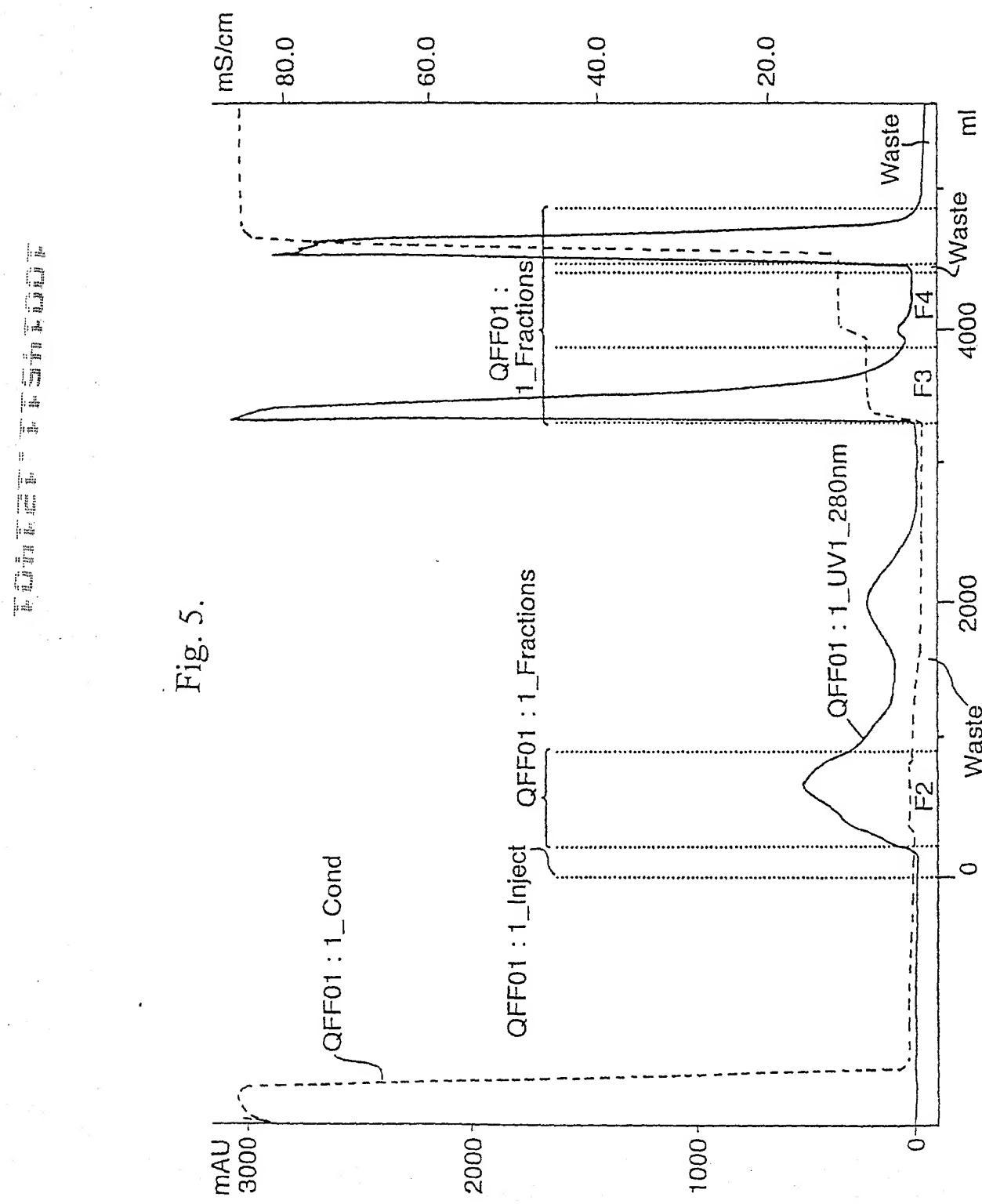


Fig. 6.

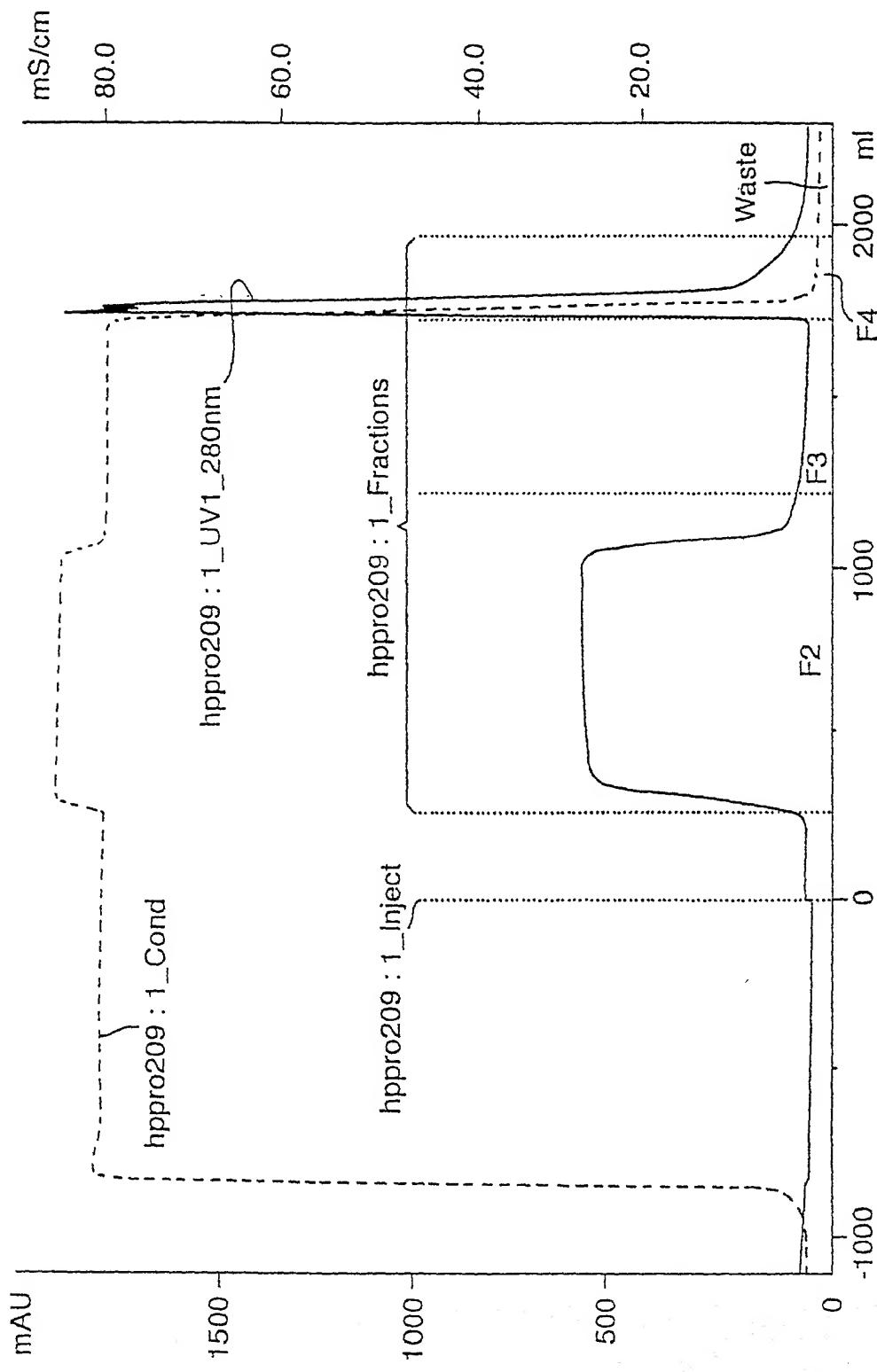


Fig. 7.

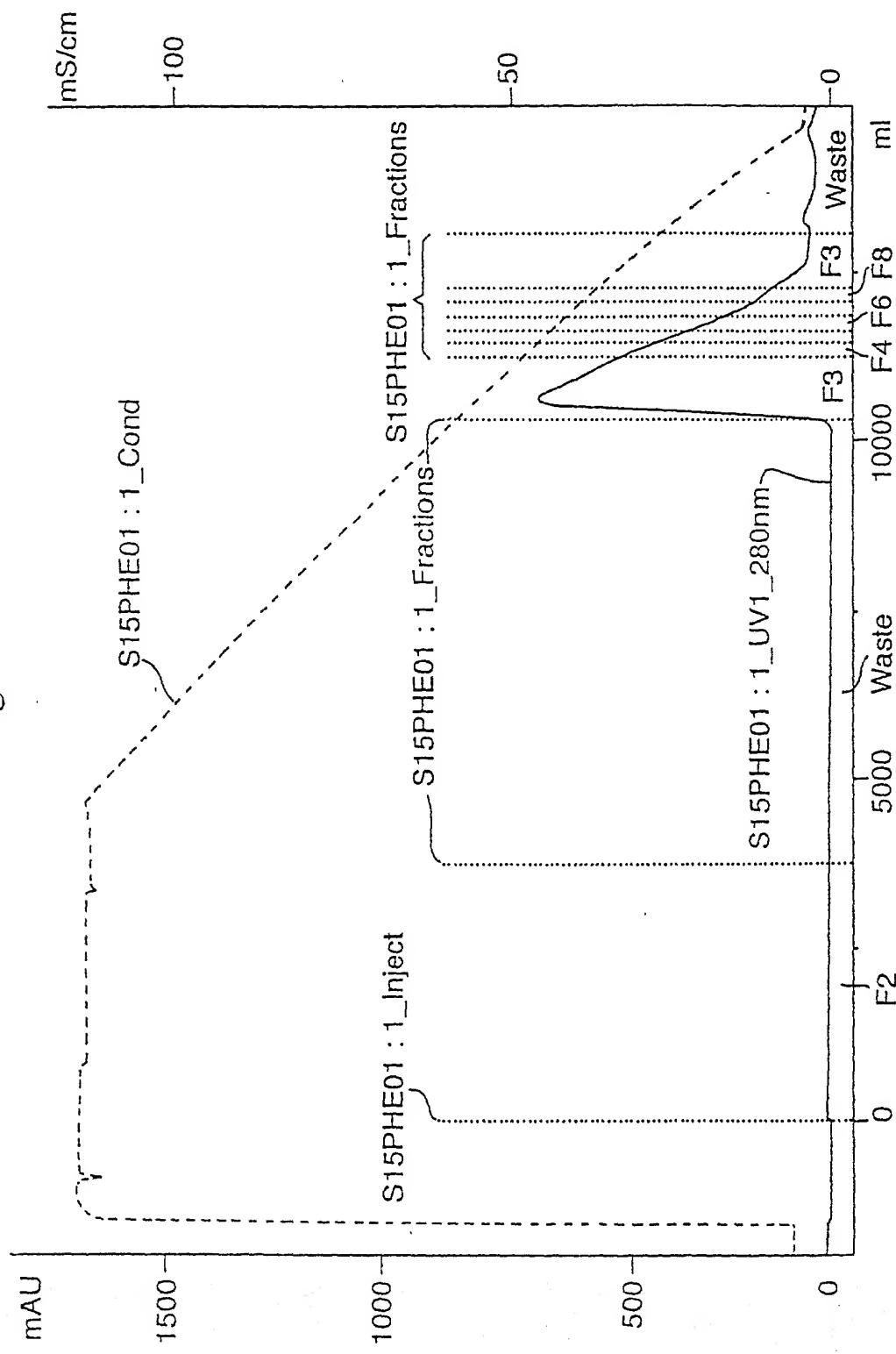


Fig. 8.

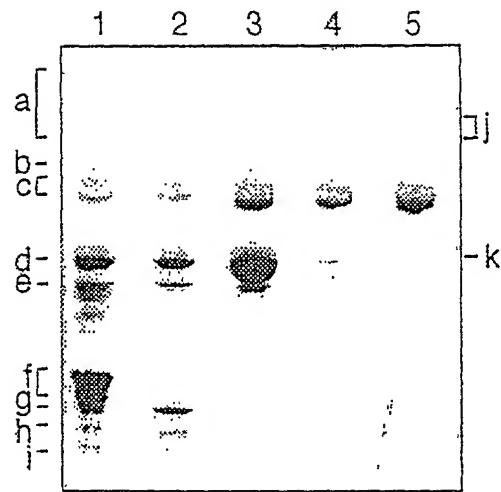


Fig. 9.

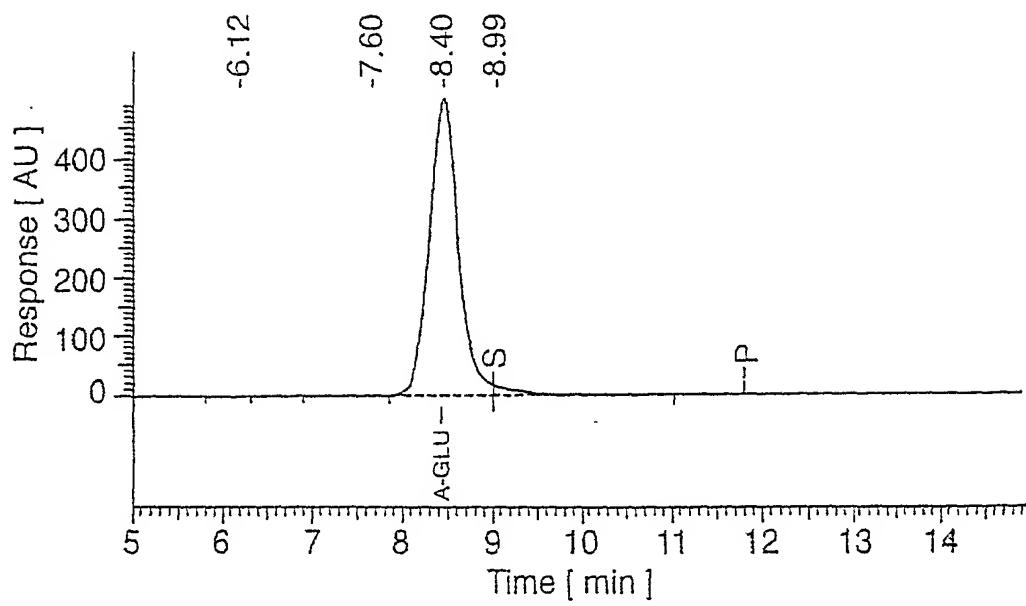


Fig. 10.

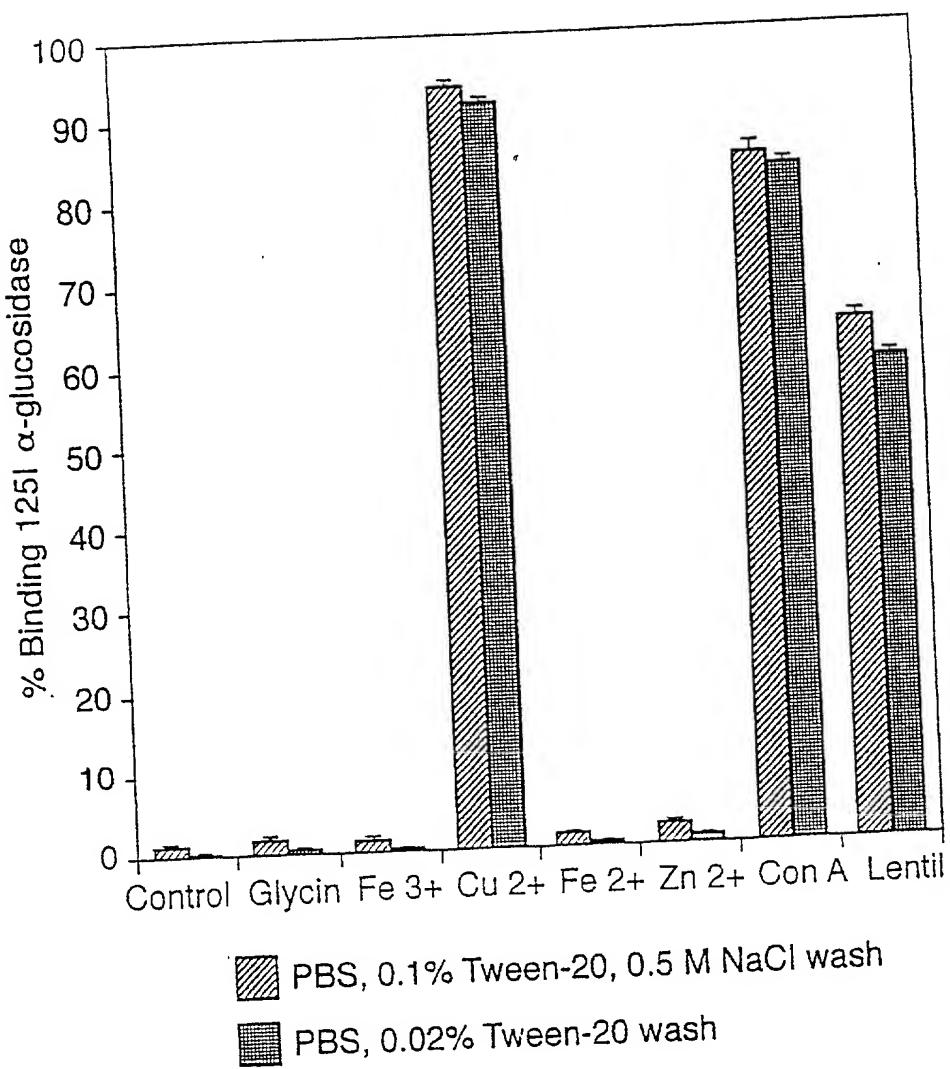


Fig. 11. A.

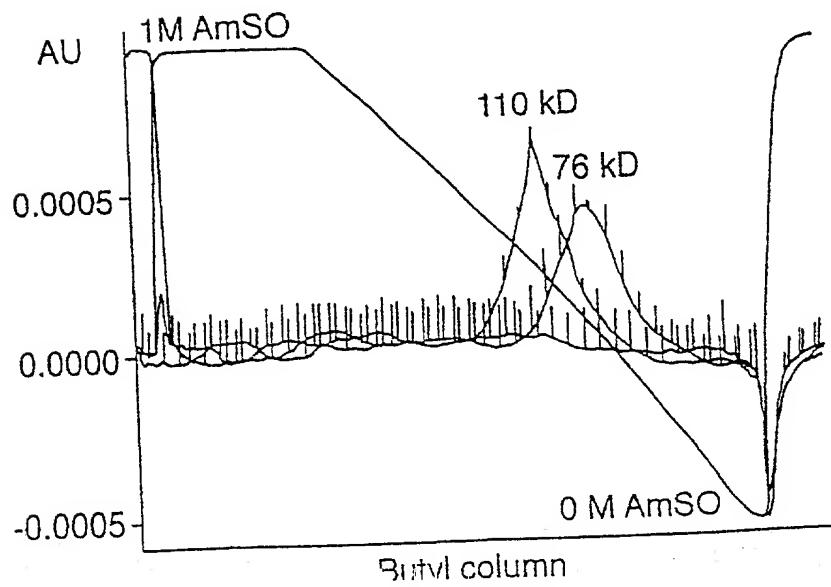


Fig. 11. B.

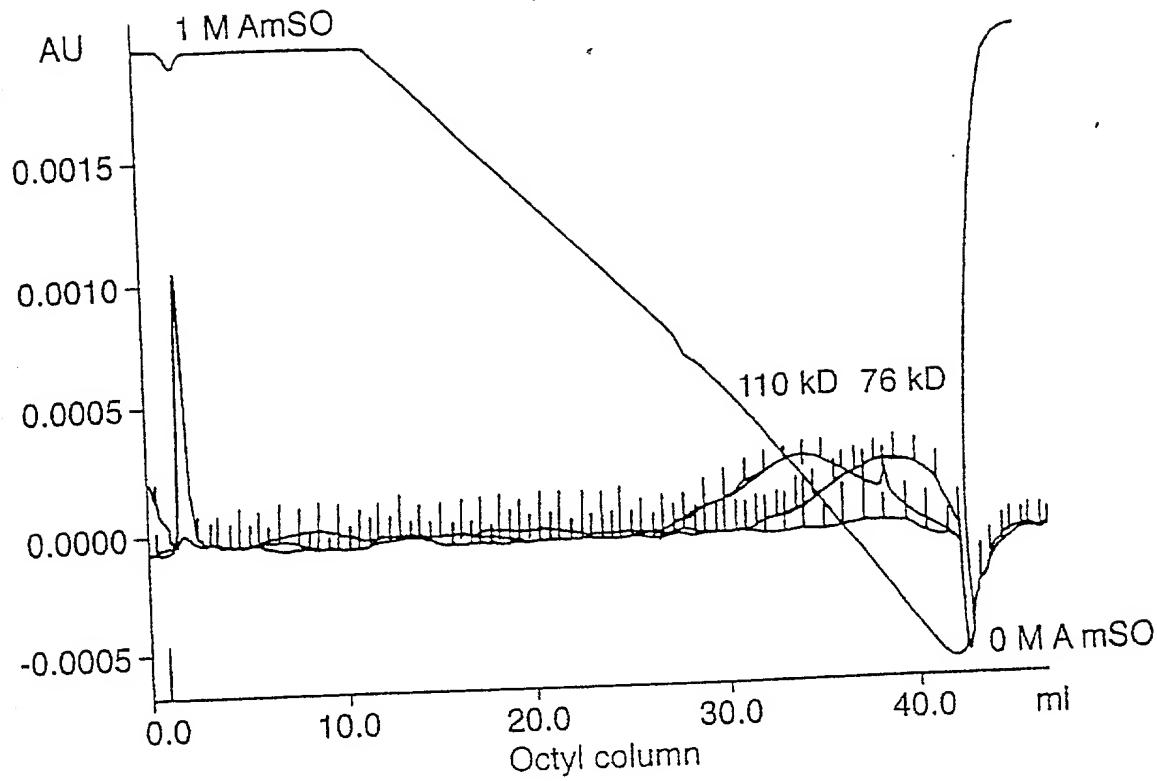


Fig. 11. C.

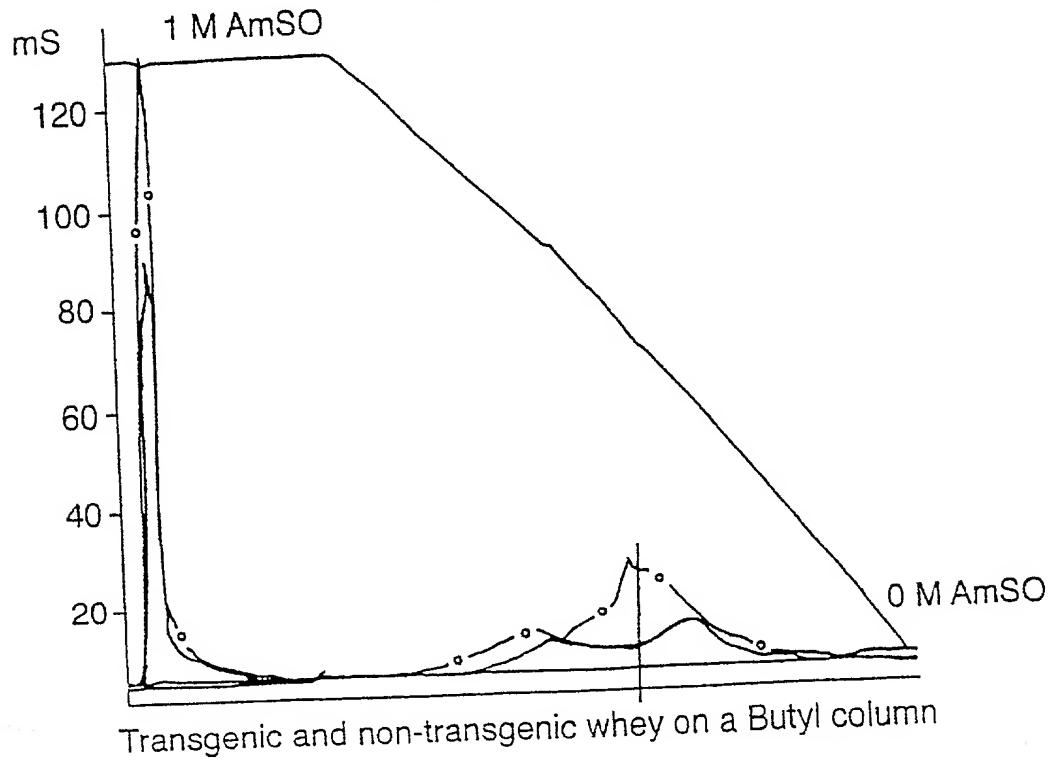


Fig. 11. D.

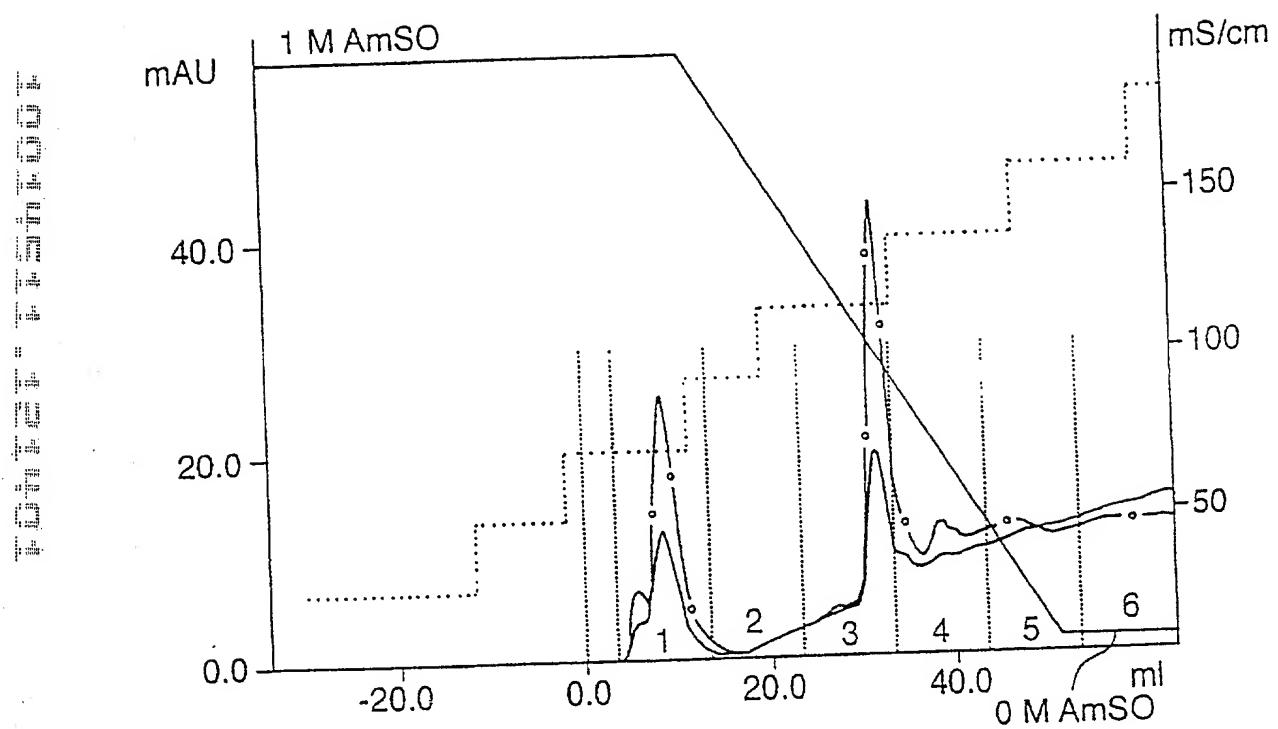


Fig. 12.

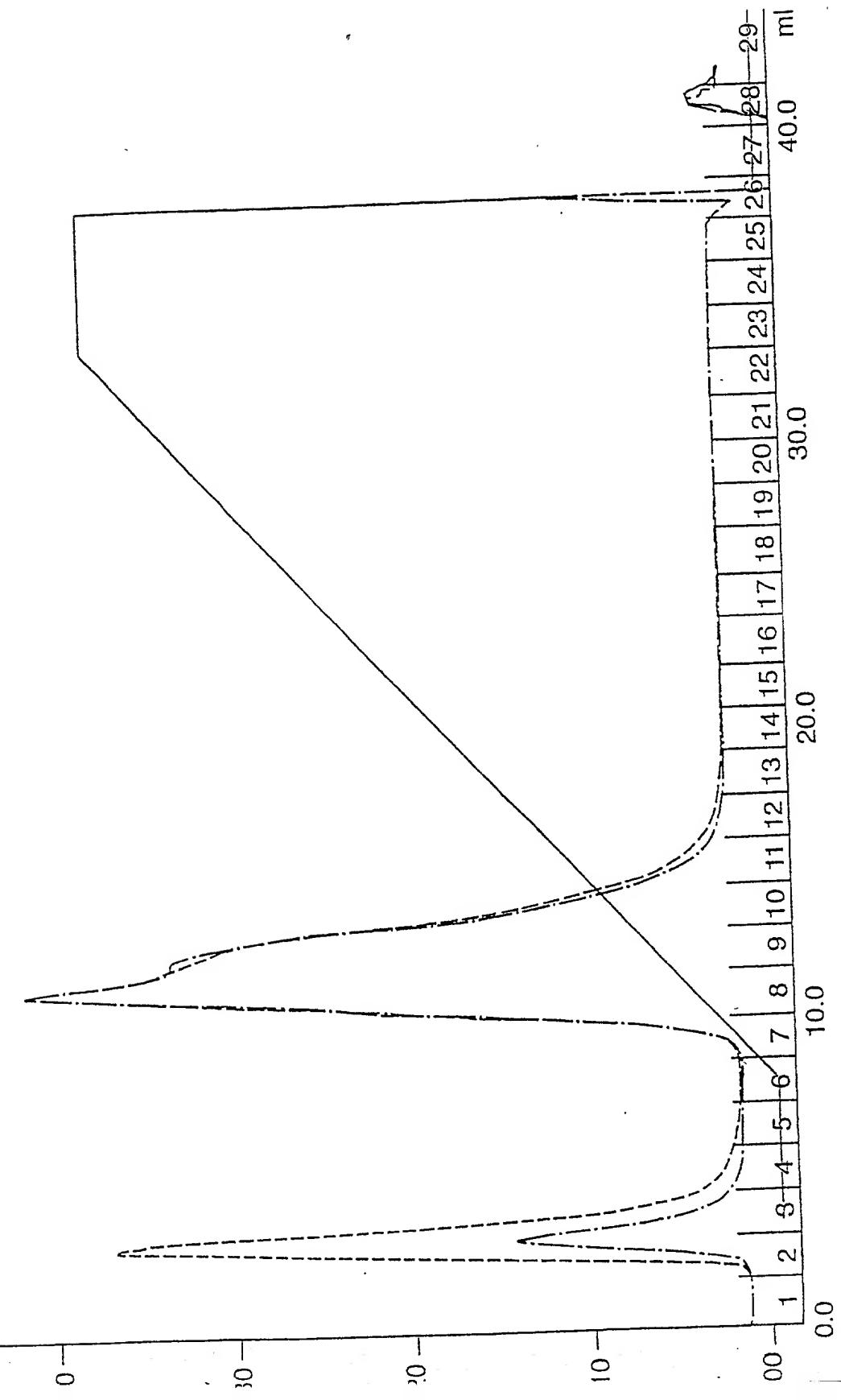


Fig. 13. A.
transgenic whey

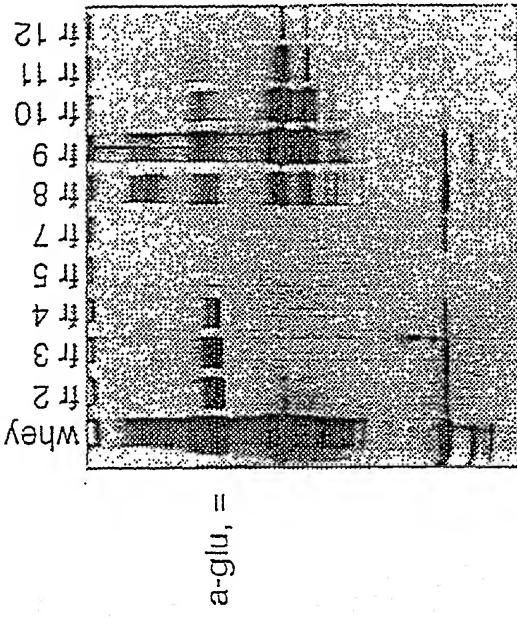


Fig. 13. B.
non-transgenic whey

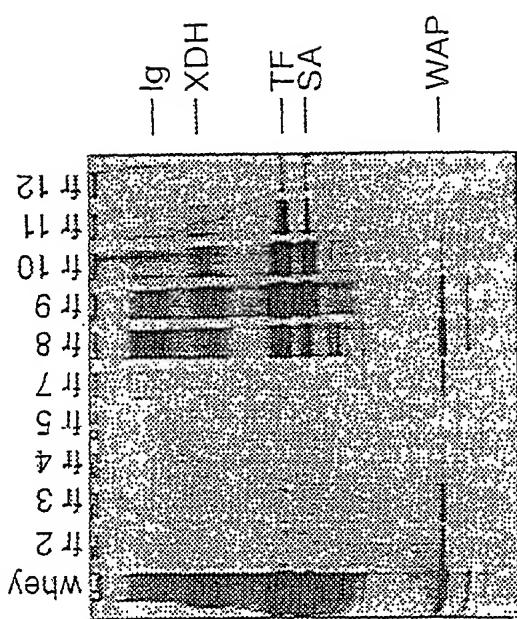
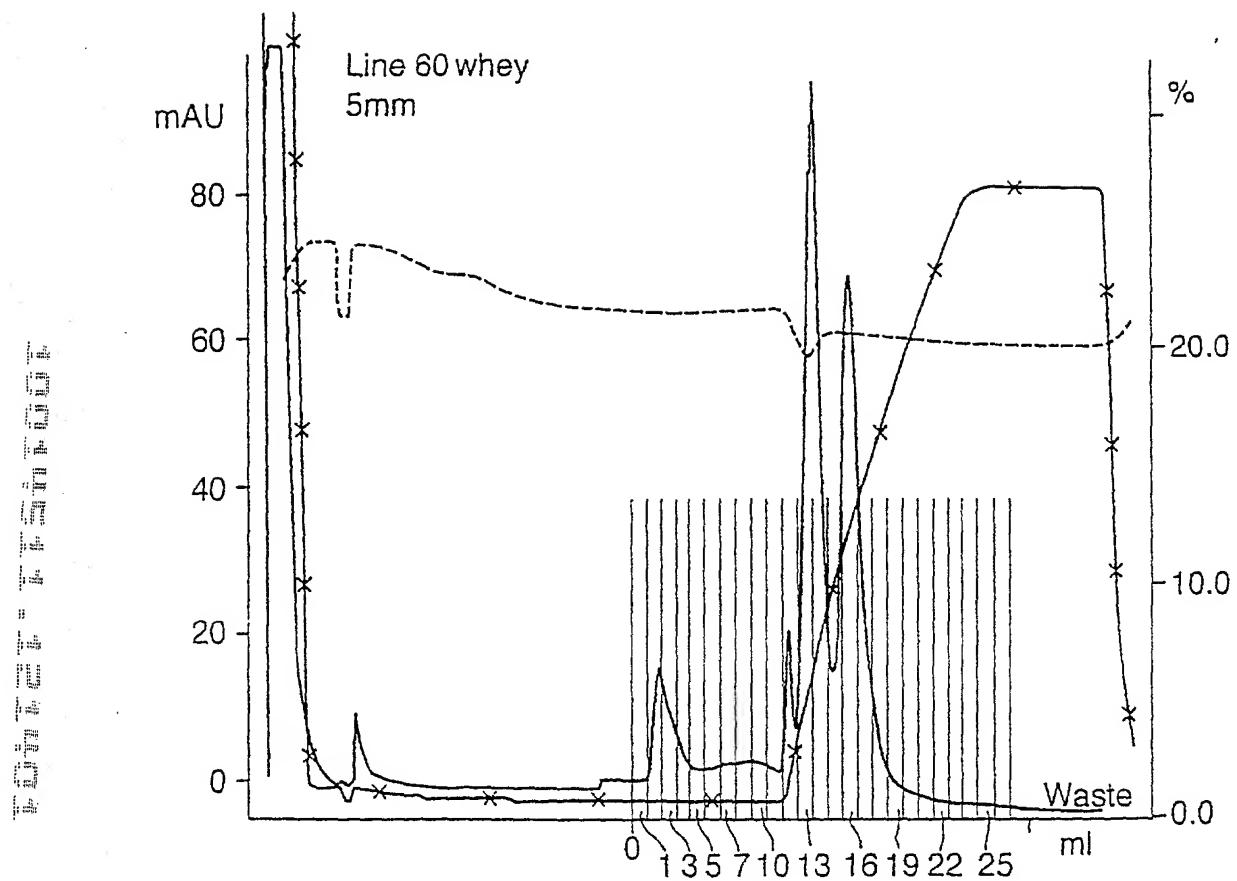


Fig. 14.



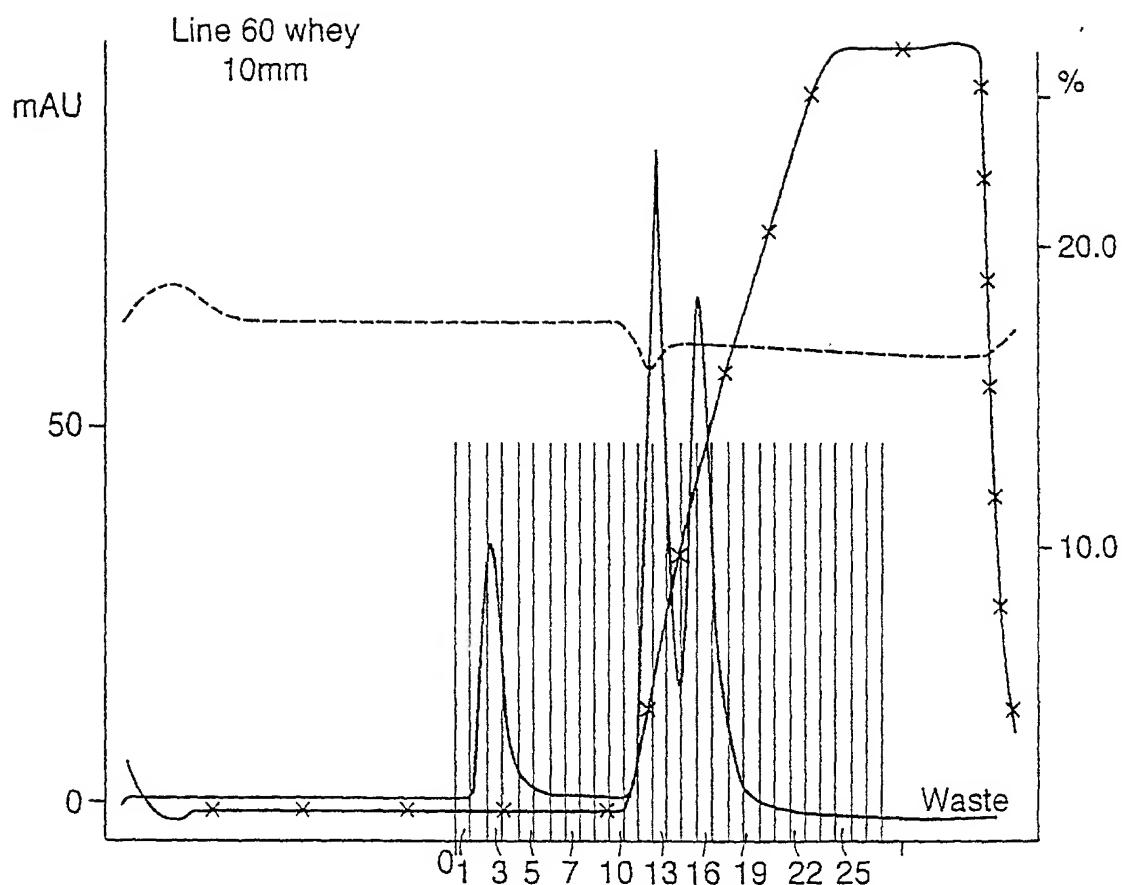
12099801:1_UV1_280nm

12099801:1_pH

12099801:1_Cond%

12099801:1_Fractions

Fig. 15.



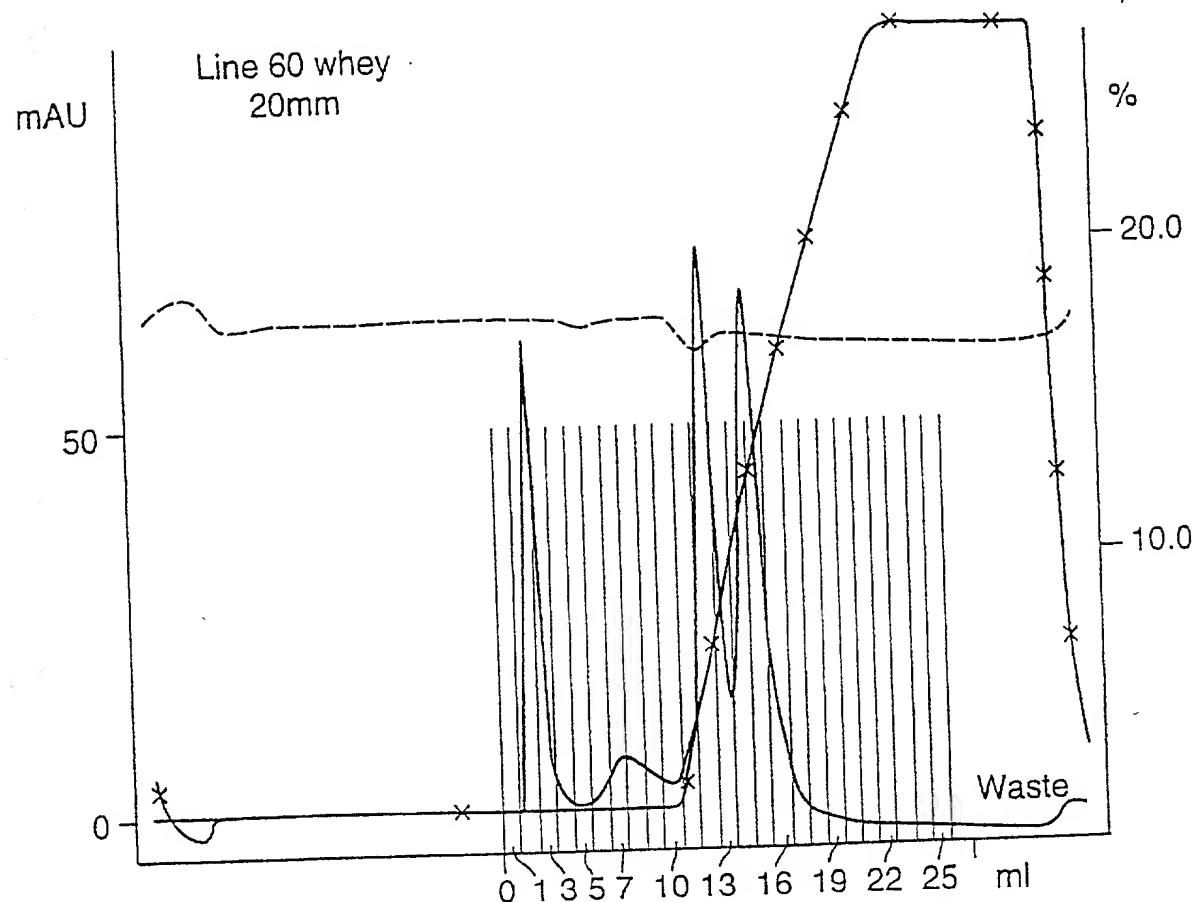
12099802:11_UV1_280nm

12099802:11_pH

12099802:11_Cond%

12099802:11_Fractions

Fig. 16.



12099803:12_UV1_280nm

12099803:12_pH

* * * 12099803:12_Cond%

12099803:12_Fractions

Fig. 17.

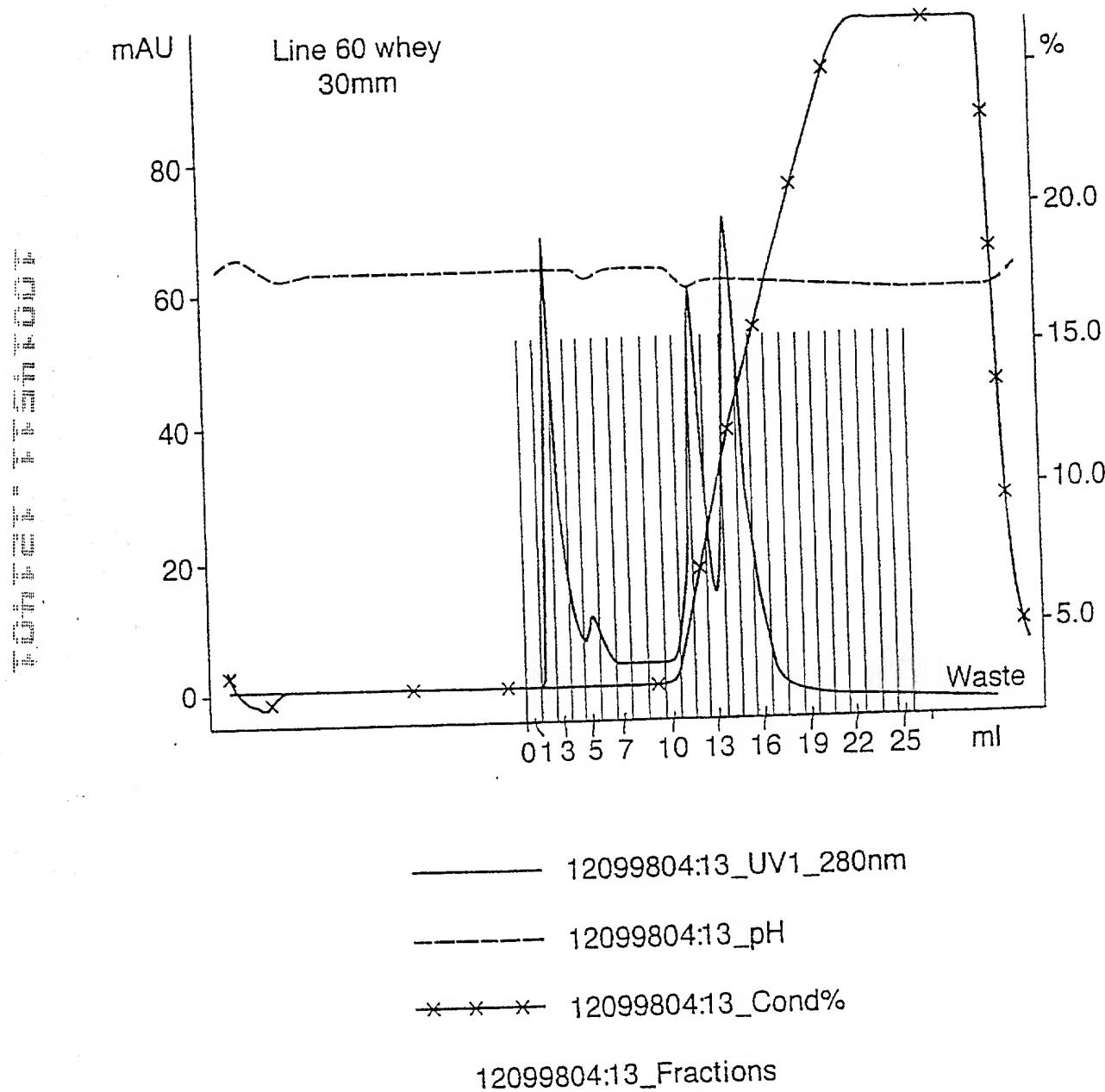
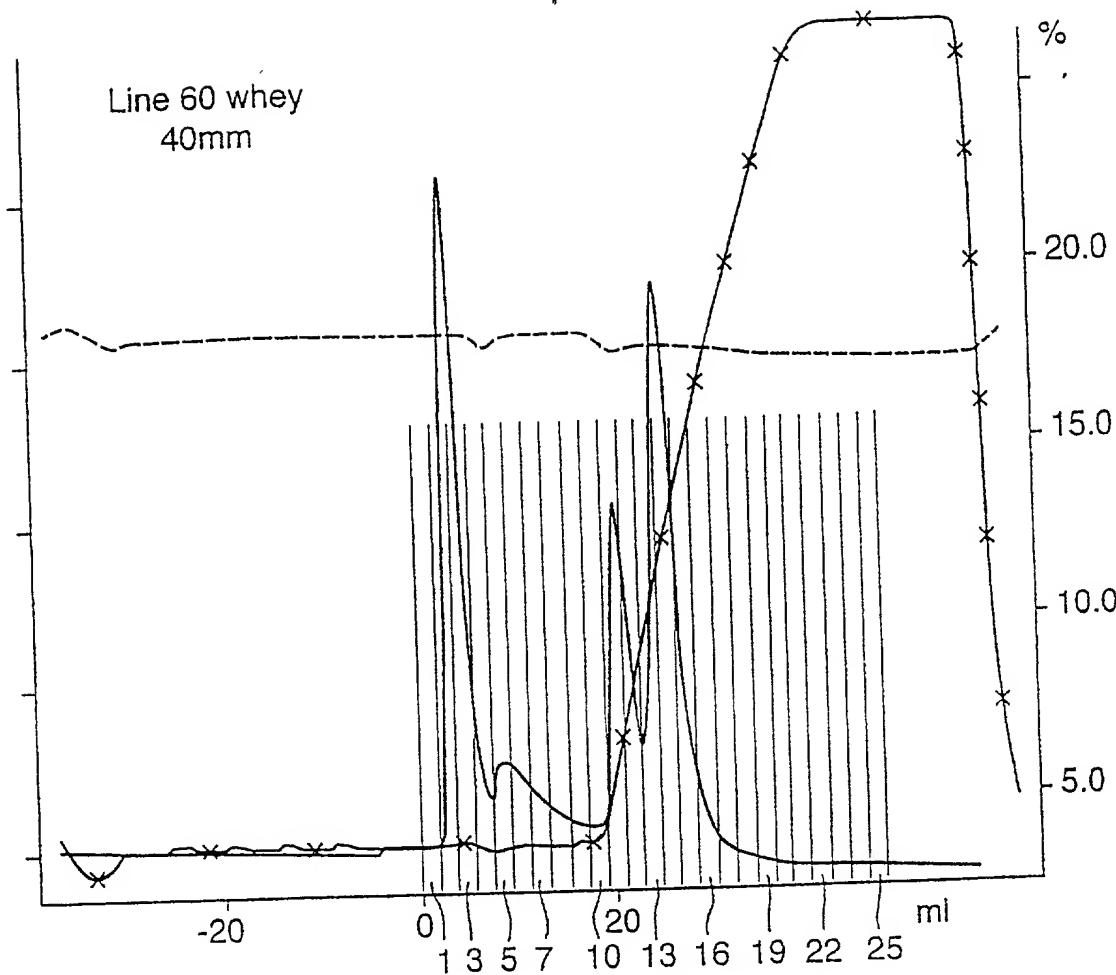


Fig. 18.



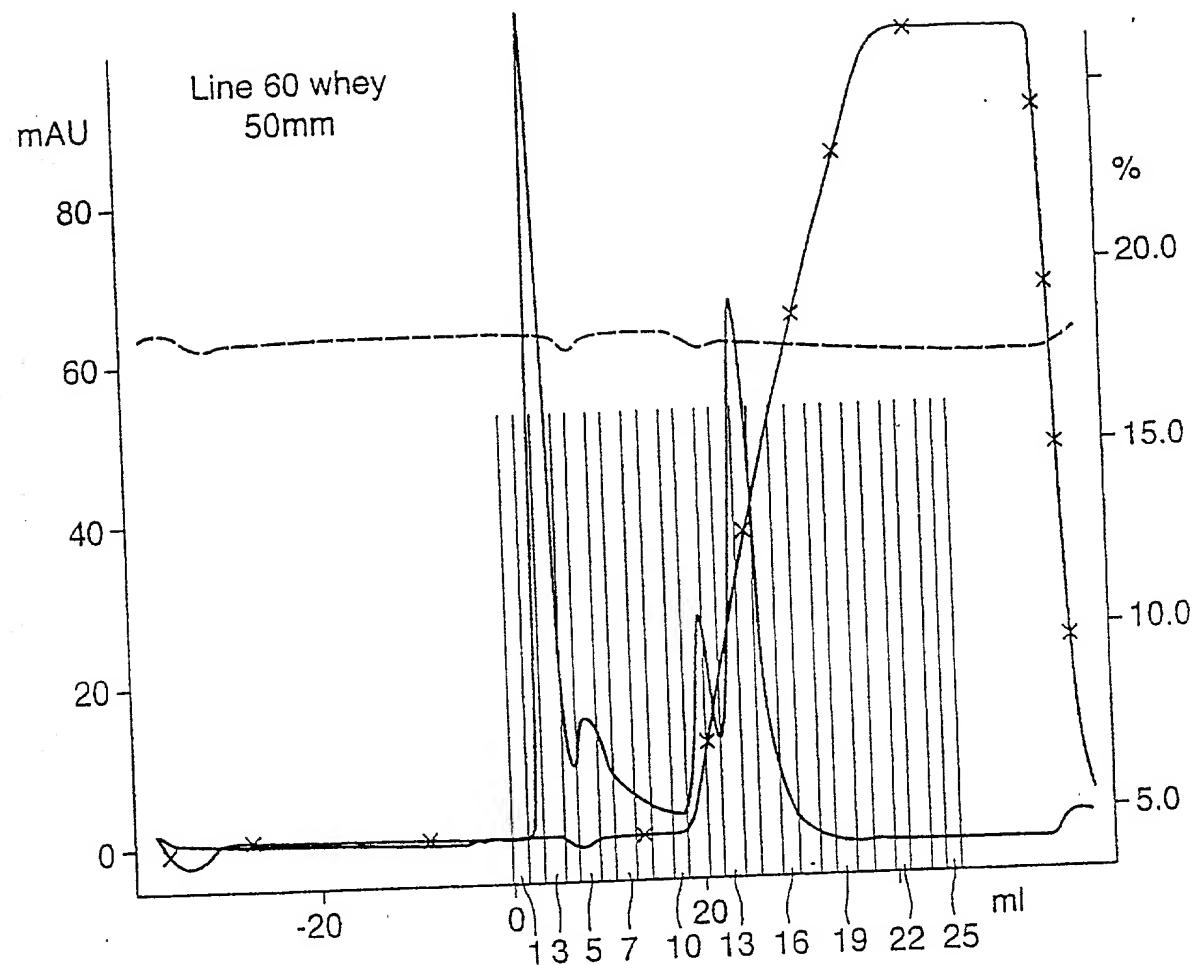
— 121099805:1_UV1_280nm

- - - 121099805:1_pH

——*—* 121099805:1_Cond%

121099805:1_Fractions

Fig. 19.



121099806:1_UV1_280nm

121099806:1_pH

121099806:1_Cond%

121099806:1_Fractions

Fig. 20.

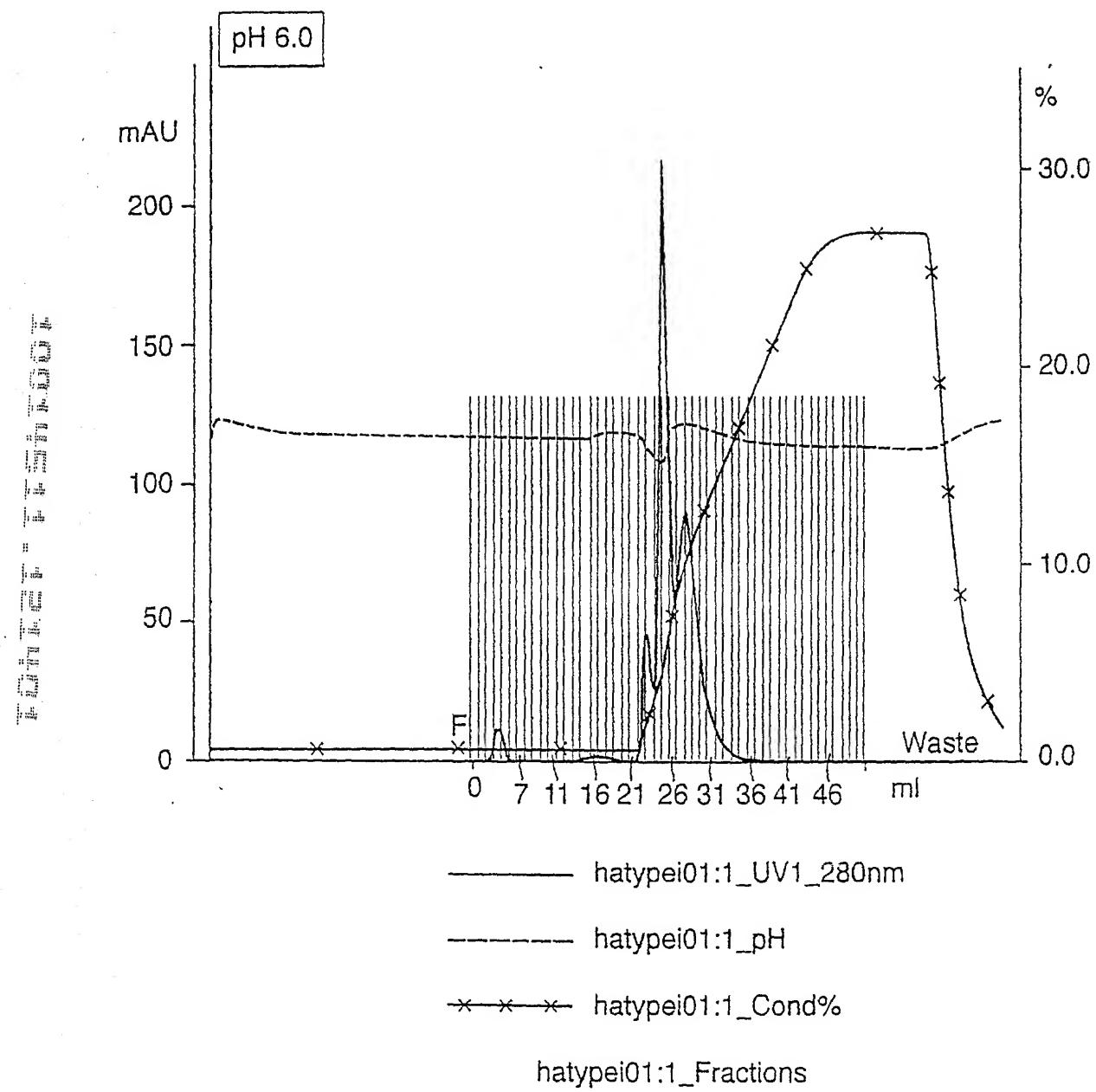


Fig. 21.

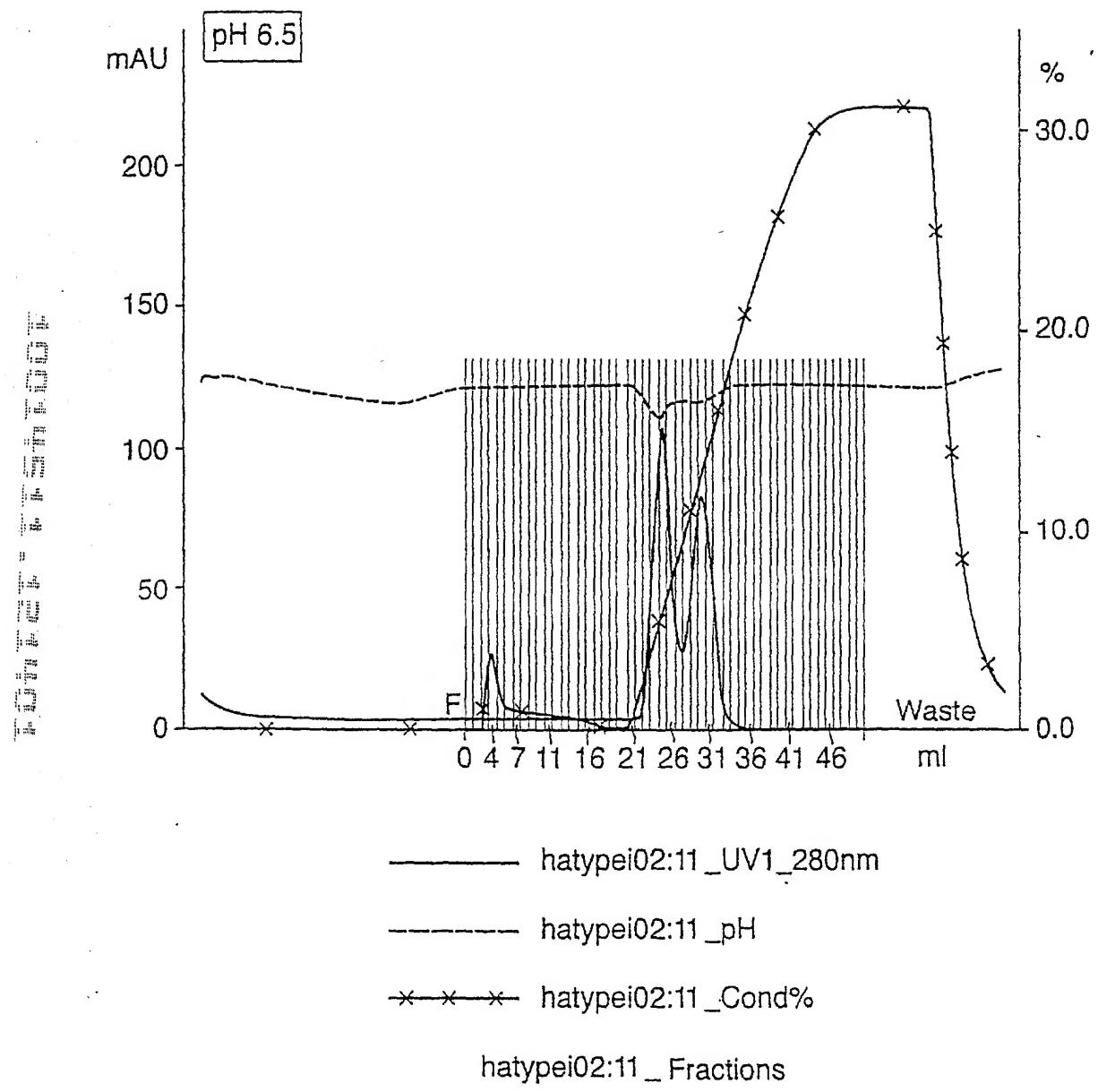


Fig. 22.

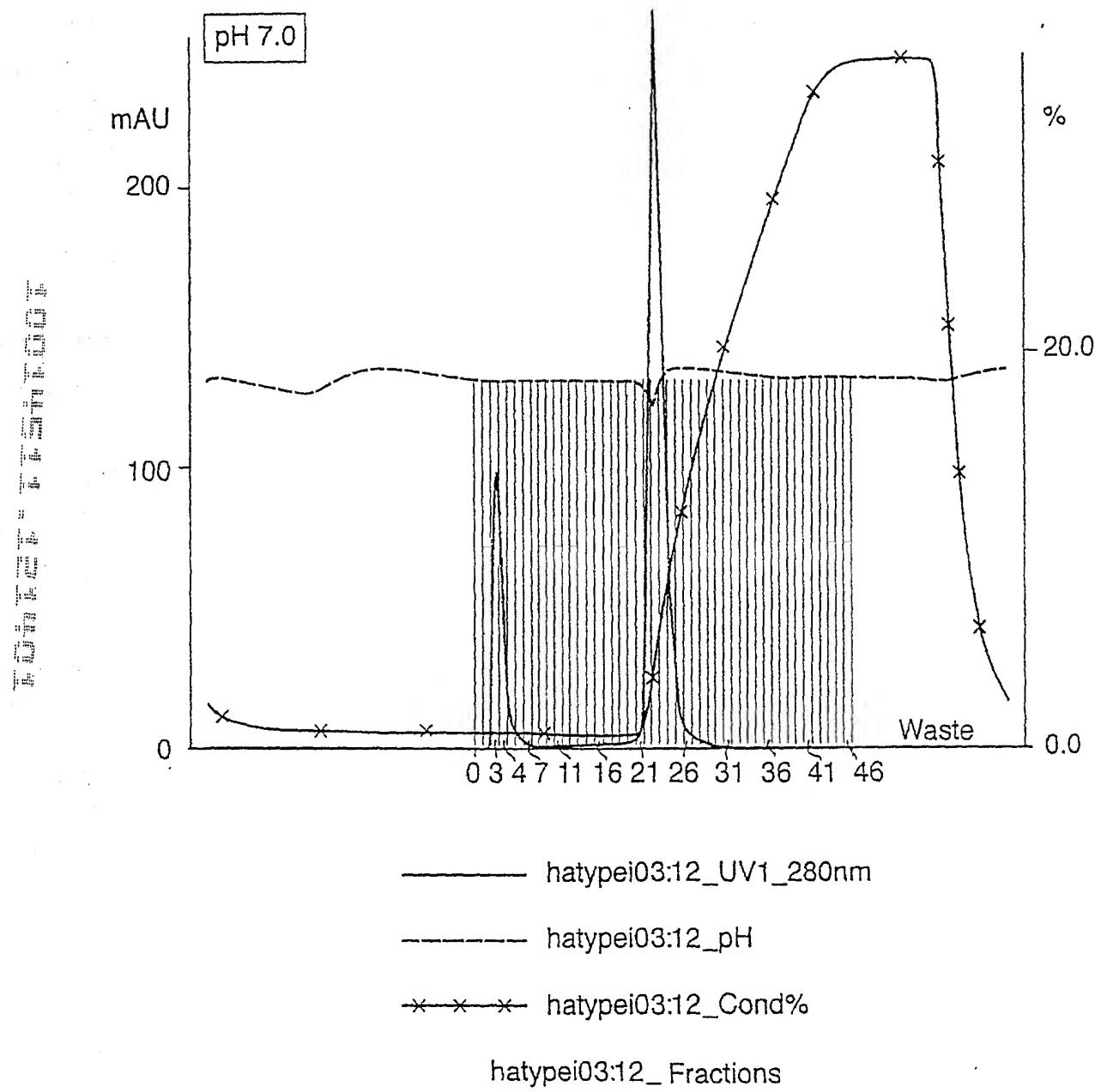


Fig. 23.

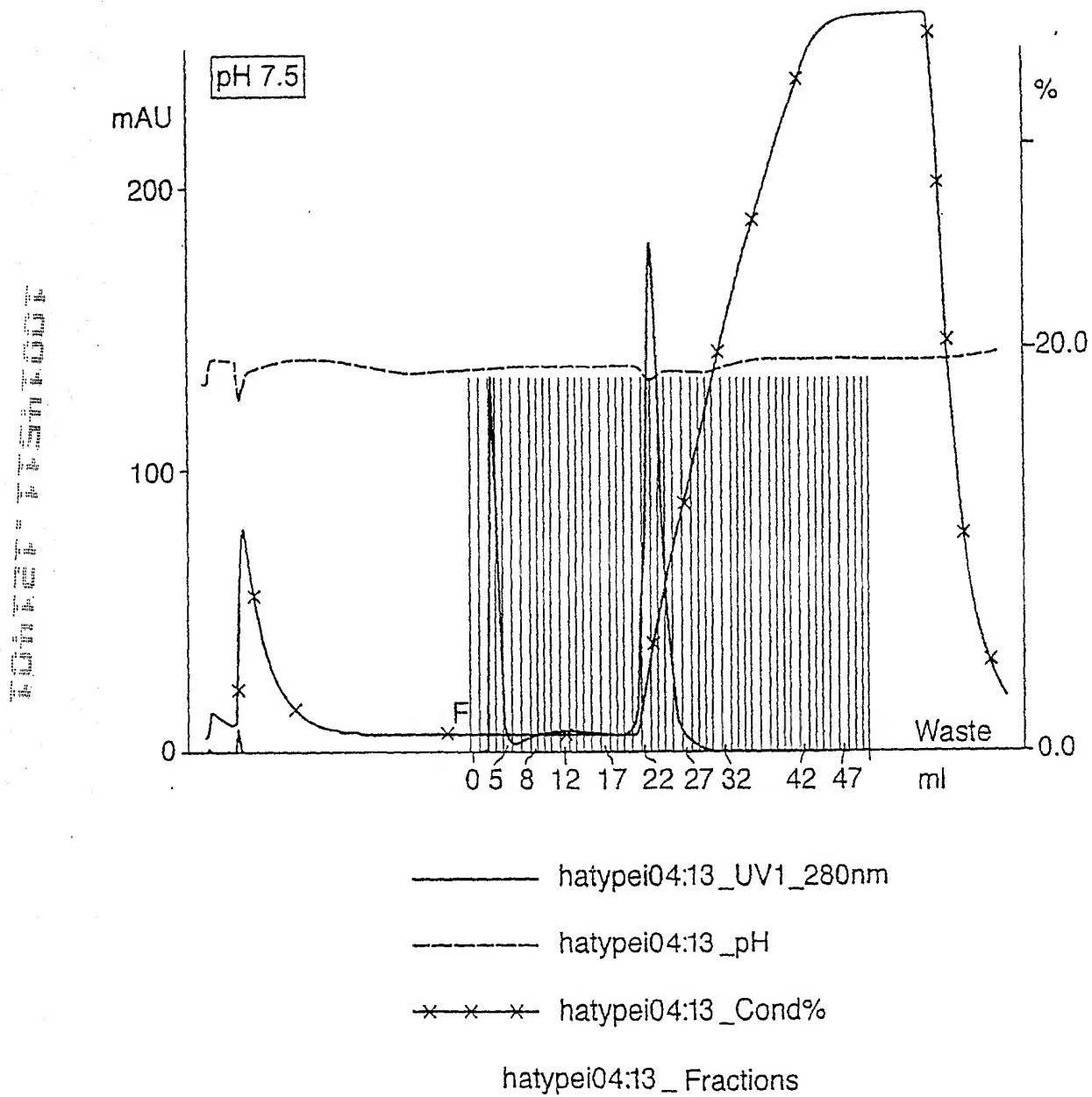


Fig. 24.

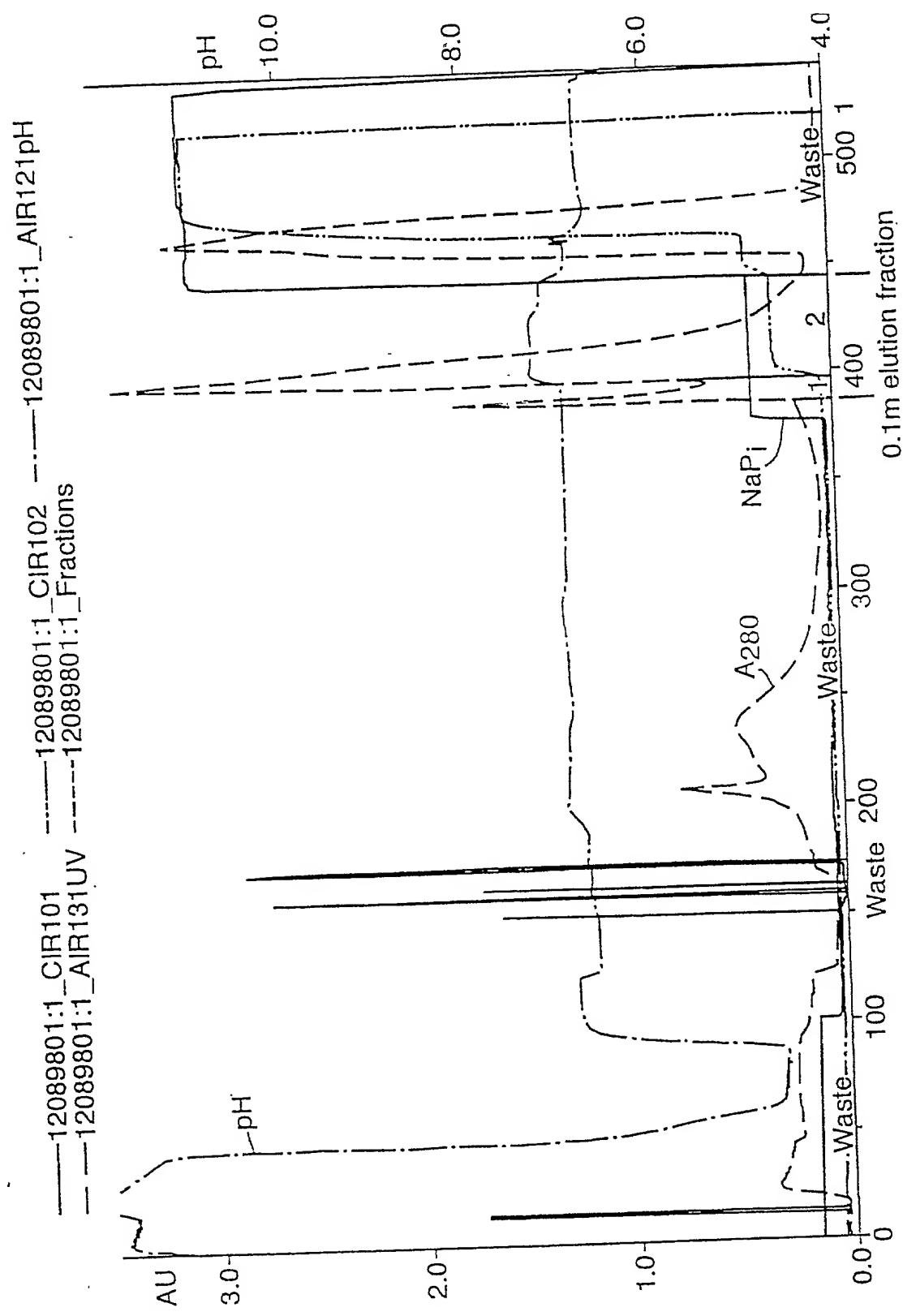


Fig. 25.

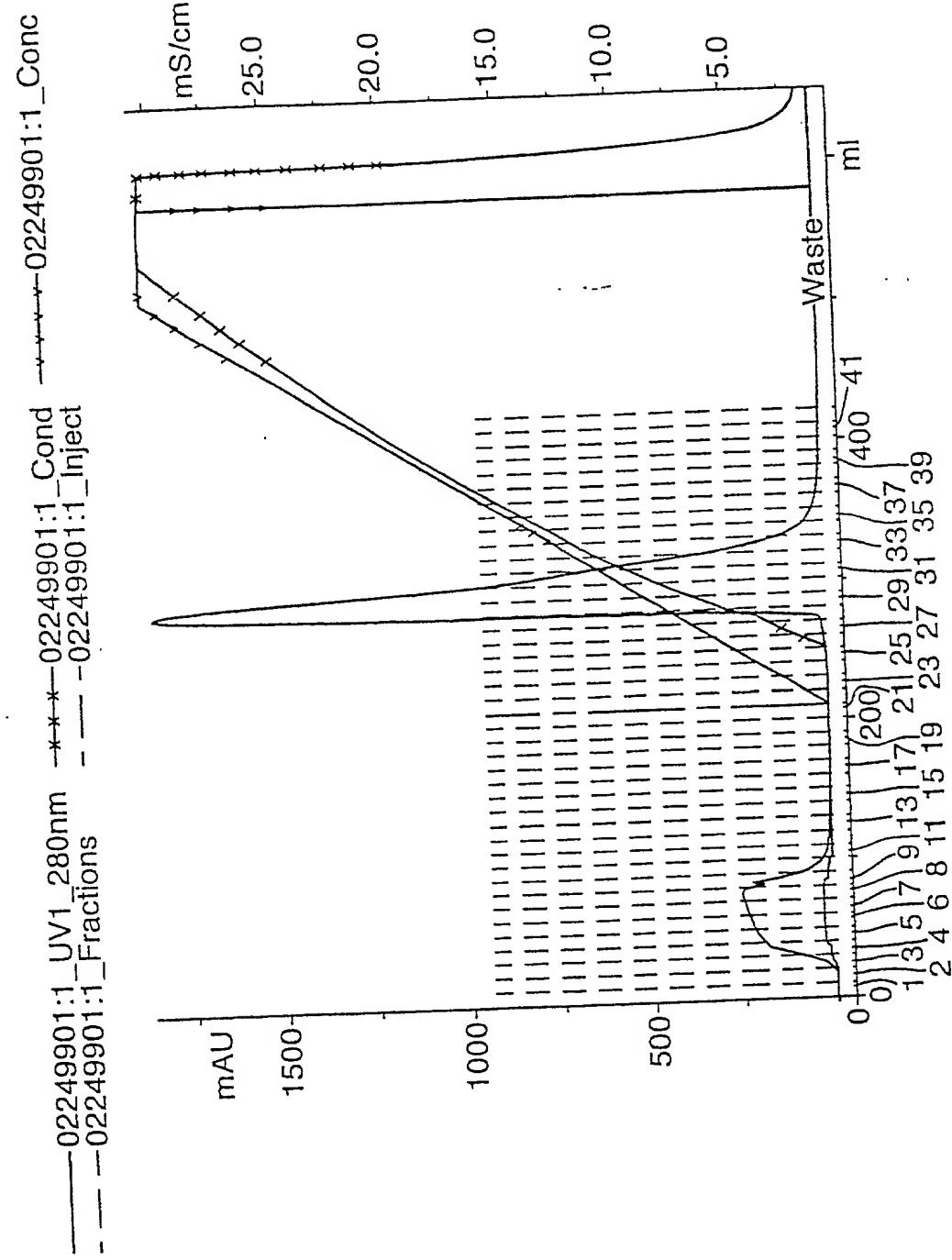


Fig. 26.

XK16/15 80°C
cHT type I 10mM Napi pH 6.5 ; QFF eluate
Run 02249901/02259901/02269901

